

Express Mail No.: EL 501 640 114 US

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Application of: Pramod K. Srivastava

Application No.: To be assigned

Group Art Unit: To be assigned

Filed: Concurrently herewith

Examiner: To be assigned

For: COMPLEXES OF ALPHA (2)
MACROGLOBULIN AND ANTIGENIC
MOLECULES FOR IMMUNOTHERAPY

Attorney Docket No.: 8449-178-999

TRANSMITTAL OF SEQUENCE LISTING UNDER 37 C.F.R. § 1.821

Assistant Commissioner for Patents
Washington, D.C. 20231

Sir:

In accordance with 37 C.F.R. § 1.821, Applicant submits herewith a Sequence Listing in paper and computer-readable form pursuant to 37 C.F.R. §§ 1.821(c) and (e).

I hereby state that the content of the paper and computer-readable copies of the Sequence Listing, submitted in accordance with 37 C.F.R. §§ 1.821(c) and (e), respectively, are the same.

Respectfully submitted,

Date: June 4, 2001

Adriane M. Antler 32,605
Adriane M. Antler (Reg. No.)

By: Eileen E. Falvey 46,097
Eileen E. Falvey (Reg. No.)

PENNIE & EDMONDS LLP
1155 Avenue of the Americas
New York, New York 10036-2711
(212) 790-9090

SEQUENCE LISTING

<110> Srivstava, Pramod K

<120> COMPLEXES OF ALPHA (2) MACROGLOBULIN AND ANTIGENIC MOLECULES FOR IMMUNOTHERAPY

<130> 8449-178

<150> 09/625,139

<151> 2000-07-25

<150> 60/209,266

<151> 2000-06-02

<160> 5

<170> FastSEQ for Windows Version 3.0

<210> 1

<211> 14849

<212> DNA

<213> Mus musculus

<400> 1

cgctgctccc	cgccagtgca	ctgaggaggc	ggaaacgggg	gagcccctag	tgctccatca	60
ggcccctacc	aaggcacccc	catcggggtcc	acgcccccca	ccccccaccc	cgccctcctcc	120
caattgtgca	tttttgagc	cggagtcggc	tccgagatgg	ggctgtgagc	ttcgccctgg	180
gagggggaga	ggagcgagga	gtaaagcagg	ggtgaagggt	tcgaatttgg	gggcaggggg	240
cgcacccgcg	tcagcaggcc	cttcccaggg	ggctcggaac	tgtaccattt	cacctatgcc	300
cctggttcgc	tttgcttaag	gaaggataag	atagaagagt	cggggagagg	aagataaagg	360
gggaccccc	aattgggggg	ggcgaggaca	agaagtaaca	ggaccagagg	gtgggggctg	420
ctgtttgcat	cggcccacac	catgctgacc	ccgccgttgc	tgctgctcgt	gccgctgctt	480
tcagctctgg	tctccggggc	cactatggat	gcccctaaaa	cttgcagccc	taagcagttt	540
gcctgcagag	accaaatac	ctgtatctca	aagggtctgg	ggtgtgacgg	tgaaagagat	600
tgccccgacg	gctctgatga	agcccctgag	atctgtccac	agagtaaagc	ccagagatgc	660
ccgccaaatg	agcacagttg	tctggggact	gagctatgtg	tccccatgtc	tcgtctctgc	720
aacgggatcc	aggactgcat	ggatggctca	gacgagggtg	ctcactgccc	agagctccga	780
gccaaactgtt	ctcgaatggg	ttgtcaacac	cattgtgtac	ctacacccag	tggggcccacg	840
tgctactgta	acagcagctt	ccagctcgag	gcagatggca	agacgtgcaa	agattttgac	900
gagtgttcgg	tgtatggcac	ctgcagccag	ctttgcacca	acacagatgg	ctccttcaca	960
tgtggctgtg	ttgaaggcta	cctgctgcaa	ccggacaacc	gctcctgcaa	ggccaagaat	1020
gagccagtag	atcgcccgcc	agtgtactg	attgccaaact	ctcagaacat	cctagctacg	1080
tacctgagtg	gggcccaggt	gtctaccatc	acaccaccca	gcacccgaca	aaccacggcc	1140
atggacttca	gttatgccaa	tgagaccgta	tgctgggtgc	acgttgggga	cagtgtctgcc	1200
cagacacagc	tcaagtgtgc	ccggatgcct	ggcctgaagg	gctttgtgga	tgagcatacc	1260
atcaacatct	ccctcagcct	gcaccacgtg	gagcagatgg	caatcgactg	gctgacggga	1320
aactttctact	ttgtcgacga	cattgacgac	aggatctttg	tctgtaaccg	aaacggggac	1380
acctgtgtca	ctctgctgga	cctggaactc	tacaaccca	aaggcatcgc	cttggaaccc	1440
gccatgggga	aggtgttctt	cactgactac	gggcagatcc	caaagggtgga	gcgctgtgac	1500
atggatggac	agaaccgcac	caagctgggtg	gatagcaaga	tcgtgtttcc	acacggcatc	1560
accctggacc	tggtcagccg	cctcgtctac	tgggcgggacg	cctacctaga	ctacatcgag	1620
gtggtagact	acgaagggaa	gggtcggcag	accatcatcc	aaggcatcct	gatcgagcac	1680
ctgtacggcc	tgaccgtgtt	tgagaactat	ctctacgcca	ccaactcgga	caatgccaac	1740
acgcagcaga	agacgagcgt	gatccgagtg	aaccggttca	acagtactga	gtaccaggtc	1800
gtcaccctgtg	tggacaaggg	tggtgccctg	catatctacc	accagcgacg	ccagccccga	1860
gtgcggagtc	acgcctgtga	gaatgaccag	tacgggaagc	caggtggctg	ctccgacatc	1920

gtctctgcag	acttgcccaa	cgcccacggg	ctggctgtgg	actgggtctc	ccgaaatctg	5460
ttttggacaa	gttacgacac	caacaagaag	cagattaacg	tggcccggct	ggacggctcc	5520
ttcaagaatg	cggtggtgca	gggcctggag	cagccccacg	gcctggtcgt	ccacccgctt	5580
cgtggccaag	tctactggac	tgatggggac	aacatcagca	tggccaacat	ggatgggagc	5640
aaccacactc	tgctcttcag	tggccagaag	ggccctgtgg	ggttggccat	tgacttcctt	5700
gagagcaaac	tctactggat	cagctctggg	aaccacacaa	tcaaccgttg	caatctggat	5760
gggagcgagc	tggaggtcat	cgacaccatg	cggagccagc	tgggcaaggc	cactgccctg	5820
gccatcatgg	gggacaagct	gtggtgggca	gatcaggtgt	cagagaagat	gggcacgtgc	5880
aacaaagccg	atggctctgg	gtccgtggtg	ctgcggaaca	gtaccacgtt	ggttatgcac	5940
atgaagggtg	atgacgagag	catccagcta	gagcatgagg	gcaccaaccc	ctgcagtgtc	6000
aacaacggag	actgttccca	gctctgcctg	ccaacatcag	agacgactcg	ctcctgtatg	6060
tgtacagccg	gttacagcct	cgggagcgga	cagcaggcct	gtgagggtgt	gggctctttt	6120
ctcctgtact	ctgtacatga	gggaattcgg	gggtattccac	tagatcccaa	tgacaagtcg	6180
gatgccctgg	tcccagtgtc	cggaacttca	ctggctgtcg	gaatcgactt	ccatgccgaa	6240
aatgacacta	tttattgggt	ggatatgggc	ctaagcacca	tcagcagggc	caagcgtgac	6300
cagacatggc	gagaggatgt	ggtgaccaac	ggtattggcc	gtgtggaggg	catcgccgtg	6360
gactggatcg	caggcaacat	atactggacg	gaccagggct	tcgatgtcat	cgaggttgcc	6420
cggctcaatg	gctcttttcg	ttatgtggtc	atttcccagg	gtctggacaa	gcctcgggcc	6480
atcactgtcc	acccagagaa	ggggactctg	ttctggaccg	agtgggggtca	ttaccacagt	6540
attgagcggg	ctcgccttga	tggcacagag	agagtgggtg	tggttaatgt	cagcatcagc	6600
tggcccaatg	gcatctcagt	agactatcag	ggcggcaagc	tctactgggtg	tgatgctcgg	6660
atggacaaga	tcgagcgcct	cgacactgga	acgggcgaga	accgggaggt	ggtcctgtcc	6720
agcaataaca	tggatatggt	ctccgtgtcc	gtgtttgagg	acttcatcta	ctggagtgc	6780
agaactcacg	ccaatggctc	catcaagcgc	ggctgcaaa	acaatgctac	agactccgtg	6840
cctctgagga	caggcattgg	tgttcagctt	aaagacatca	aggtcttcaa	cagggacagg	6900
cagaagggta	ccaatgtgtg	cgcggtagcc	aacggcgggt	gccagcagct	ctgcttgat	6960
cggggtggcg	gacagcgagc	ctgtgcctgt	gccacggga	tgctggcaga	agacggggcc	7020
tcattgccgag	agtacgctgg	ctacctgtct	tactcagagc	ggaccatcct	caagagcatc	7080
caactgtcgg	atgagcgtaa	cctcaacgca	ccggtgcagc	cctttgaaga	ccccgagcac	7140
atgaaaaatg	tcattgcctc	ggcctttgac	taccgagcag	gcacctcccc	ggggacccct	7200
aaccgcatct	tcttcagtga	catccacttt	gggaacatcc	agcagatcaa	tgacgatggc	7260
tcgggcagga	ccaccatcgt	ggaaaatgtg	ggctctgtgg	aaggcctggc	ctatcacctg	7320
ggctgggaca	cactgtactg	gacaagctac	accacatcca	ccatcacccg	ccacaccgtg	7380
gaccagactc	gccagggggc	cttcgagagg	gagacagtca	tcaccatgtc	cggagacgac	7440
cacccgagag	cctttgtgct	ggatgagtgc	cagaaactga	tgttctggac	caattggaac	7500
gagctccatc	caagcatcat	gcgggcagcc	ctatccggag	ccaacgtcct	gaccctcatt	7560
gagaaggaca	tccgcacgcc	caatgggttg	gccatcgacc	accgggcgga	gaagctgtac	7620
ttctcggatg	ccaccttgga	caagatcgag	cgctgcgagt	acgacggctc	ccaccgctat	7680
gtgatcctaa	agtcggagcc	cgtccacccc	tttgggttgg	cgggtgtacg	agagcacatt	7740
ttctggactg	actgggtgcg	gcgggctgtg	cagcagagcca	acaagtatgt	gggcagcgac	7800
atgaagctgc	ttcgggtgga	cattccccag	caacccatgg	gcattcatcg	cgtggccaat	7860
gacaccaaca	gctgtgaact	ctccccctgc	cgtatcaaca	atggaggctg	ccaggatctg	7920
tgtctgctca	cccaccaagg	ccacgtcaac	tgttcctgtc	gagggggccg	gatcctccag	7980
gaggacttca	cctgccgggg	tgtgaactcc	tcttgtcggg	cacaagatga	gtttgagtgt	8040
gccaatgggg	aatgtatcag	cttcagcctc	acctgtgatg	gcgtctccca	ctgcaaggac	8100
aagtccgatg	agaagccctc	ctactgcaac	tcacgcgctc	gcaagaagac	tttccgccag	8160
tgtaacaatg	gccgctgtgt	atccaacatg	ctgtggtgca	atgggggtgga	ttactgtggg	8220
gatggctctg	atgagatacc	ttgcaacaag	actgcctgtg	gtgtgggtga	gttccgctgc	8280
cgggatgggt	cctgcatcgg	gaactccagt	cgtgcgaacc	agtttgtgga	ttgtgaggat	8340
gcctcggatg	agatgaattg	cagtgccaca	gactgcagca	gctattttccg	cctgggctgtg	8400
aaaggtgtcc	tcttccagcc	gtgcgagcgg	acatccctgt	gctacgcacc	tagctgggtg	8460
tgtgatggcg	ccaacgactg	tggagactac	agcgatgaac	gtgactgtcc	aggtgtgaag	8520
cgccttaggt	gcccgtctaa	ttacttttgc	tgccccagcg	ggcgtgtgat	ccccatgagc	8580
tggacgtgtg	acaaggagga	tgactgtgag	aacggcgagg	atgagaccca	ctgcaacaag	8640
ttctgctcag	aggcacagtt	cgagtgccag	aaccaccggt	gtatctccaa	gcagtggctg	8700
tgtgacggta	gcgatgattg	cggggatggc	tccgatgagg	cagctcactg	tgaaggcaag	8760
acatgtggcc	cctcctcctt	ctcctgtccc	ggcaccacag	tgtgtgtccc	tgagcgctgg	8820
ctctgtgatg	gcgacaagga	ctgtaccgat	ggcgcggtg	agagtgtcac	tgctggctgc	8880

ctgtacaaca	gcacctgtga	tgaccgtgag	ttcatgtgcc	agaaccgctt	gtgtattccc	8940
aagcatttcg	tgtgcgacca	tgaccgtgac	tgtgctgatg	gctctgatga	atcccctgag	9000
tgtgagtagc	caacctgctg	gcccattgaa	ttccgctgtg	ccaatgggcg	ttgtctgagc	9060
tcccgtcagt	gggaatgtga	tggggagaat	gactgtcacg	accacagcga	tgaggctccc	9120
aagaaccac	actgcaccag	cccagagcac	aaatgcaatg	cctcatcaca	gttcctgtgc	9180
agcagcgggc	gctgctgtgg	tgaggcgttg	ctctgcaacg	gccaggacga	ctgtggggac	9240
ggttcagacg	aacgcgggtg	ccatgtcaac	gagtgtctca	gccgcaagct	cagtggctgc	9300
agtcaggact	gcgaggacct	caagataggg	tttaagtgcc	gctgtcgccc	gggcttccgg	9360
ctaaaggacg	atggcaggac	ctgtgcccac	ctggatgagt	gcagcaccac	cttcccctgc	9420
agccagctct	gcataaacac	ccacggaagt	tacaagtgtc	tgtgtgtgga	gggctatgca	9480
ccccgtggcg	gtgaccccc	cagctgcaaa	gctgtgaccg	atgaggagcc	atttctcatc	9540
tttgccaacc	ggtactacct	gcggaagctc	aacctggacg	gctccaacta	cacactgctt	9600
aagcagggcg	tgaacaatgc	ggtcgccttg	gcatttgact	accgagagca	gatgatctac	9660
tggacgggcg	tgaccaccca	gggcagcatg	attcgcagga	tgcacctcaa	cggcagcaac	9720
gtgcagggtt	tgcaccggac	gggccttagt	aaccagatg	ggctcgctgt	ggactgggtg	9780
gggtggcaacc	tgtactgggt	tgacaagggc	agagatacca	ttgagggtgt	caagcttaac	9840
ggggcctatc	ggacagtgtc	ggtcagctct	ggcctccggg	agcccagagc	tctggtagt	9900
gatgtacaga	atgggtacct	gtactggaca	gactgggggt	accactcact	gatcggccgg	9960
attggcatgg	atggatctgg	ccgcagcatc	atcgtggaca	ctaagatcac	atggcccaat	10020
ggcctgaccg	tggactacgt	cacggaacgc	atctactggg	ctgacgcccg	tgaggactac	10080
atcgagttcg	ccagcctgga	tggctccaac	cgtcacgttg	tgctgagcca	agacatccca	10140
cacatctttg	cgctgacctt	at ttgaagac	tacgtctact	ggacagactg	ggaaacgaag	10200
tccatcaaac	gggcccacaa	gaccacgggt	gccaaacaaa	cactcctcat	cagcaccctg	10260
caccggccca	tggacttaca	tgtattccac	gccctgcgcc	agccagatgt	gcccatacac	10320
ccctgcaaa	tcaacaatgg	tggctgcagc	aacctgtgcc	tgctgtcccc	tggggggtggt	10380
cacaagtgcg	cctgccccac	caacttctat	ctgggtggcg	atggccgtac	ctgtgtgtcc	10440
aactgcacag	caagccagtt	tgtgtgcaaa	aatgacaagt	gcatacccct	ctggtggaag	10500
tgtgacacgg	aggacgactg	tggggatcac	tcagacgagc	ctccagactg	tcccagagtt	10560
aagtgccggc	caggccagtt	ccagtgtctc	accggcatct	gcaccaaccc	tgcttctatc	10620
tgtgatgggg	acaatgactg	ccaagacaat	agtgcagagg	ccaattgcga	cattcacgtc	10680
tgcttgccca	gccaatcaaa	gtgcaccaac	accaaccgct	gcatttctgg	catcttccgt	10740
tgcaatgggc	aggacaactg	cggggacggc	gaggatgagc	gggattgccc	tgaggtgacc	10800
tgcccccaca	accagttcca	gtgtccatc	accaagcgct	gcataccctc	cgctctgggt	10860
tgtgacaggg	ataatcactg	tgtggacggc	agtgatgagc	ctgccaaact	tacccaaatg	10920
acctgtggag	tggatgagtt	ccgtgcaag	gatttctggc	gctgcatccc	cgcgcgctgg	10980
aagtgtgacg	gagaagatga	ctgtggggat	ggttcagatg	agcccaagga	agagtgtgat	11040
gagcgacct	gtgagccata	ccagttccgc	tgcaaaaaca	accgctgtgt	cccaggccgt	11100
tggcaatgtg	actacgacaa	cgactgcgga	gataactcgg	acgaggagag	ctgcacacct	11160
cggccctgct	ctgagagtga	gtttttctgt	gccaatggcc	gctgcatcgc	tgggcgctgg	11220
aagtgtgatg	gggaccatga	ctgtgcccac	ggctcagacg	agaaagactg	cacccccgcg	11280
tgtgatatgg	accagttcca	gtgcaagagt	ggccactgca	tccccctgcg	ctggccgtgt	11340
gacgcggatg	ctgactgtat	ggacggcagt	gacgaggaag	cctgtggcac	tggggtgagg	11400
acctgcccac	tggatgagtt	tcaatgtaac	aacaccttgt	gcaagccgct	ggcctggaag	11460
tgtgatggag	aggacgactg	tggggacaac	tcagatgaga	accccgagga	atgcgcccgg	11520
ttcatctgcc	ctcccaaccg	gcctttccgc	tgcaagaatg	accgagtctg	cctgtggatt	11580
gggcgcagct	gtgatggcgt	ggacaactgt	ggagatggga	ctgacgagga	ggactgtgag	11640
ccccccacgg	cccagaaccc	ccactgcaaa	gacaagaagg	agttcctgtg	ccgaaaccag	11700
cgctgtctat	catcctccct	gcgtgtgaac	atgttcgatg	actgcggcga	tggctccgat	11760
gaagaagatt	gcagcatcga	ccccaaagct	accagctgtg	ccaccaatgc	cagcatgtgt	11820
ggggacgaag	ctcgttgtgt	gcgcactgag	aaagctgcct	actgtgcctg	ccgctcgggc	11880
ttccatactg	tgccggggcca	gcccggatgc	caggacatca	acgagtgcct	gcgctttggt	11940
acctgctctc	agctctggaa	caaaccacaa	ggaggccacc	tctgcagctg	tgcccgaac	12000
ttcatgaaga	cacacaacac	ctgcaaaagt	gaaggctccg	agtaccaggt	gctatacatc	12060
gcggatgaca	acgagatccg	cagcttgttc	ccggggccacc	cccactcagc	ctacgagcag	12120
acattccagg	gcgatgagag	tgtccgcata	gatgccatgg	atgtccatgt	caaggccggc	12180
cgtgtctact	ggactaaactg	gcacacgggc	acaatctcct	acaggagcct	gccccctgcc	12240
gcccctccta	ccacttccaa	ccgccaccgg	aggcagatcg	accgggggtg	caccacctc	12300
aatatttcag	ggctgaagat	gccgaggggt	atcgctatcg	actgggtggc	cgggaatgtg	12360

tactggaccg	attccggccg	agacgtgatt	gaggtggcgc	aaatgaaggg	cgagaaccgc	12420
aagacgctca	tctcgggcat	gattgatgag	ccccatgcc	tcgtggtgga	ccctctgagg	12480
ggcaccatgt	actggtcaga	ctgggggaac	cacccaaga	ttgaaacagc	agcgatggat	12540
ggcacccttc	gggagactct	cgtgcaagac	aacattcagt	ggcctacagg	gctggctgtg	12600
gactatcaca	atgaacggct	ctactgggca	gatgccaagc	tttcggtcac	cggcagcatc	12660
cggctcaacg	gcaactgacc	cattgtggct	gctgacagca	aacgaggcct	aagtcacccc	12720
ttcagcatcg	atgtgtttga	agactacatc	tacggagtca	cttacatcaa	taatcgtgtc	12780
ttcaagatcc	acaagtttgg	acacagcccc	ttgtacaacc	taactggggg	cctgagccat	12840
gcctctgatg	tagtccttta	ccatcaacac	aagcagcctg	aagtgaccaa	cccctgtgac	12900
cgcaagaaat	gcgaatggct	gtgtctgctg	agccccagcg	ggcctgtctg	cacctgtccc	12960
aatggaaaga	ggctggataa	tggcacctgt	gtgcctgtgc	cctctccaac	accccccca	13020
gatgccccca	ggcctgggaa	ctgcactctg	cagtgcctca	atgggtggtg	ttgtttcctc	13080
aacgctcggg	ggcagcccaa	gtgccgttgc	cagccccgtt	acacaggcga	taagtgtgag	13140
ctggatcagt	gctgggaata	ctgtcacaac	ggaggcacct	gtgcggcttc	cccatctggc	13200
atgcccacgt	gocgtgtcc	cactggcttc	acgggcccc	aatgcaccgc	acaggtgtgt	13260
gcaggctact	gctctaaca	cagcacctgc	accgtcaacc	agggcaacca	gccccagtgc	13320
cgatgtctac	ctggcttcc	gggcgaccgt	tgccagtacc	ggcagtgtc	tggcttctgt	13380
gagaactttg	gcacctgtca	gatggctgct	gatggctccc	gacaatgtcg	ctgcaccgtc	13440
tactttgagg	gaccaaggtg	tgaggtgaac	aagtgtagtc	gctgtctcca	aggcgctgt	13500
gtggtcaata	agcagaccgg	agatgtcaca	tgcaactgca	ctgatggccg	ggtagcccc	13560
agttgtctca	cctgcactga	tcactgtagc	aatggtggct	cctgcaccat	gaacagcaag	13620
atgatgcctg	agtgccagtg	cccgcccat	atgacaggac	cccggtgcc	ggagcaggtt	13680
gttagtcagc	aacagcctgg	gcataatggc	tccatcctga	tccctctgct	gctgcttctc	13740
ctgctgcttc	tgggtgctgg	cgtgggtgtc	tggataaagc	ggcgagtccg	aggggctaag	13800
ggcttccagc	accagcggat	gaccaatggg	gccatgaatg	tggaaattgg	aaaccctacc	13860
tacaagatgt	atgaaggtgg	agagcccgat	gatgtcgggg	gcctactgga	tgctgatttt	13920
gcccttgacc	ctgacaagcc	taccaacttc	accaaccag	tgtatgccac	gctctacatg	13980
gggggcccag	gcagccgcca	ttccctggcc	agcacggacg	agaagcgaga	actgctgggc	14040
cggggacctg	aagacgagat	aggagatccc	ttggcatagg	gccctgcccc	gacggatgtc	14100
cccagaaagc	cccctgccac	atgagtcttt	caatgaacc	cctccccagc	cggcccttct	14160
ccggccctgc	cgggtgtaca	aatgtaaaaa	tgaagggaatt	actttttata	tgtgagcgag	14220
caagcgagca	agcacagtat	tatctctttg	catttccttc	ctgcctgctc	ctcagtatcc	14280
cccccatgct	gccttgaggg	ggcggggagg	gctttgtggc	tcaaaggat	gaaggagtcc	14340
acatgttccc	taccgagcat	acccctggaa	gcctggcggc	acggcctccc	caccacgcct	14400
gtgcaagaca	ctcaacgggg	ctccgtgtcc	cagctttcct	ttccttggct	ctctgggggt	14460
agttcagggg	aggtggagtc	ctctgctgac	cctgtctgga	agatttggct	ctagctgagg	14520
aaggagtctt	ttagttgagg	gaagtcaccc	caaaccacag	ctcccacttt	caggggcacc	14580
tctcagatgg	ccatgctcag	tatcccttcc	agacaggccc	tccctctctc	agcgccccct	14640
ctgtggctcc	tagggctgaa	cacattcttt	ggtaactgtc	ccccaaagcct	cccatcccc	14700
tgagggccag	gaagagtcgg	ggcacaccaa	ggaagggcaa	gcgggcagcc	ccattttggg	14760
gacgtgaacg	ttttaataat	ttttgctgaa	ttcctttaca	actaaataac	acagatattg	14820
ttataaataa	aattgtaaaa	aaaaaaaaa				14849

<210> 2

<211> 4545

<212> PRT

<213> Mus musculus

<400> 2

Met	Leu	Thr	Pro	Pro	Leu	Leu	Leu	Leu	Val	Pro	Leu	Leu	Ser	Ala	Leu
1				5					10					15	
Val	Ser	Gly	Ala	Thr	Met	Asp	Ala	Pro	Lys	Thr	Cys	Ser	Pro	Lys	Gln
			20					25					30		
Phe	Ala	Cys	Arg	Asp	Gln	Ile	Thr	Cys	Ile	Ser	Lys	Gly	Trp	Arg	Cys
		35					40					45			

Asp	Gly	Glu	Arg	Asp	Cys	Pro	Asp	Gly	Ser	Asp	Glu	Ala	Pro	Glu	Ile	50	55	60
Cys	Pro	Gln	Ser	Lys	Ala	Gln	Arg	Cys	Pro	Pro	Asn	Glu	His	Ser	Cys	65	70	75
Leu	Gly	Thr	Glu	Leu	Cys	Val	Pro	Met	Ser	Arg	Leu	Cys	Asn	Gly	Ile	85	90	95
Gln	Asp	Cys	Met	Asp	Gly	Ser	Asp	Glu	Gly	Ala	His	Cys	Arg	Glu	Leu	100	105	110
Arg	Ala	Asn	Cys	Ser	Arg	Met	Gly	Cys	Gln	His	His	Cys	Val	Pro	Thr	115	120	125
Pro	Ser	Gly	Pro	Thr	Cys	Tyr	Cys	Asn	Ser	Ser	Phe	Gln	Leu	Glu	Ala	130	135	140
Asp	Gly	Lys	Thr	Cys	Lys	Asp	Phe	Asp	Glu	Cys	Ser	Val	Tyr	Gly	Thr	145	150	155
Cys	Ser	Gln	Leu	Cys	Thr	Asn	Thr	Asp	Gly	Ser	Phe	Thr	Cys	Gly	Cys	165	170	175
Val	Glu	Gly	Tyr	Leu	Leu	Gln	Pro	Asp	Asn	Arg	Ser	Cys	Lys	Ala	Lys	180	185	190
Asn	Glu	Pro	Val	Asp	Arg	Pro	Pro	Val	Leu	Leu	Ile	Ala	Asn	Ser	Gln	195	200	205
Asn	Ile	Leu	Ala	Thr	Tyr	Leu	Ser	Gly	Ala	Gln	Val	Ser	Thr	Ile	Thr	210	215	220
Pro	Thr	Ser	Thr	Arg	Gln	Thr	Thr	Ala	Met	Asp	Phe	Ser	Tyr	Ala	Asn	225	230	235
Glu	Thr	Val	Cys	Trp	Val	His	Val	Gly	Asp	Ser	Ala	Ala	Gln	Thr	Gln	245	250	255
Leu	Lys	Cys	Ala	Arg	Met	Pro	Gly	Leu	Lys	Gly	Phe	Val	Asp	Glu	His	260	265	270
Thr	Ile	Asn	Ile	Ser	Leu	Ser	Leu	His	His	Val	Glu	Gln	Met	Ala	Ile	275	280	285
Asp	Trp	Leu	Thr	Gly	Asn	Phe	Tyr	Phe	Val	Asp	Asp	Ile	Asp	Asp	Arg	290	295	300
Ile	Phe	Val	Cys	Asn	Arg	Asn	Gly	Asp	Thr	Cys	Val	Thr	Leu	Leu	Asp	305	310	315
Leu	Glu	Leu	Tyr	Asn	Pro	Lys	Gly	Ile	Ala	Leu	Asp	Pro	Ala	Met	Gly	325	330	335
Lys	Val	Phe	Phe	Thr	Asp	Tyr	Gly	Gln	Ile	Pro	Lys	Val	Glu	Arg	Cys	340	345	350
Asp	Met	Asp	Gly	Gln	Asn	Arg	Thr	Lys	Leu	Val	Asp	Ser	Lys	Ile	Val	355	360	365
Phe	Pro	His	Gly	Ile	Thr	Leu	Asp	Leu	Val	Ser	Arg	Leu	Val	Tyr	Trp	370	375	380
Ala	Asp	Ala	Tyr	Leu	Asp	Tyr	Ile	Glu	Val	Val	Asp	Tyr	Glu	Gly	Lys	385	390	395
Gly	Arg	Gln	Thr	Ile	Ile	Gln	Gly	Ile	Leu	Ile	Glu	His	Leu	Tyr	Gly	405	410	415
Leu	Thr	Val	Phe	Glu	Asn	Tyr	Leu	Tyr	Ala	Thr	Asn	Ser	Asp	Asn	Ala	420	425	430
Asn	Thr	Gln	Lys	Thr	Ser	Val	Ile	Arg	Val	Asn	Arg	Phe	Asn	Ser		435	440	445
Thr	Glu	Tyr	Gln	Val	Val	Thr	Arg	Val	Asp	Lys	Gly	Gly	Ala	Leu	His	450	455	460
Ile	Tyr	His	Gln	Arg	Arg	Gln	Pro	Arg	Val	Arg	Ser	His	Ala	Cys	Glu	465	470	475
Asn	Asp	Gln	Tyr	Gly	Lys	Pro	Gly	Gly	Cys	Ser	Asp	Ile	Cys	Leu	Leu	485	490	495
Ala	Asn	Ser	His	Lys	Ala	Arg	Thr	Cys	Arg	Cys	Arg	Ser	Gly	Phe	Ser	500	505	510

Leu	Gly	Ser	Asp	Gly	Lys	Ser	Cys	Lys	Lys	Pro	Glu	His	Glu	Leu	Phe
		515					520					525			
Leu	Val	Tyr	Gly	Lys	Gly	Arg	Pro	Gly	Ile	Ile	Arg	Gly	Met	Asp	Met
	530					535					540				
Gly	Ala	Lys	Val	Pro	Asp	Glu	His	Met	Ile	Pro	Ile	Glu	Asn	Leu	Met
545					550					555					560
Asn	Pro	Arg	Ala	Leu	Asp	Phe	His	Ala	Glu	Thr	Gly	Phe	Ile	Tyr	Phe
			565						570					575	
Ala	Asp	Thr	Thr	Ser	Tyr	Leu	Ile	Gly	Arg	Gln	Lys	Ile	Asp	Gly	Thr
		580						585					590		
Glu	Arg	Glu	Thr	Ile	Leu	Lys	Asp	Gly	Ile	His	Asn	Val	Glu	Gly	Val
		595				600						605			
Ala	Val	Asp	Trp	Met	Gly	Asp	Asn	Leu	Tyr	Trp	Thr	Asp	Asp	Gly	Pro
	610					615					620				
Lys	Lys	Thr	Ile	Ser	Val	Ala	Arg	Leu	Glu	Lys	Ala	Ala	Gln	Thr	Arg
625					630					635					640
Lys	Thr	Leu	Ile	Glu	Gly	Lys	Met	Thr	His	Pro	Arg	Ala	Ile	Val	Val
				645					650					655	
Asp	Pro	Leu	Asn	Gly	Trp	Met	Tyr	Trp	Thr	Asp	Trp	Glu	Glu	Asp	Pro
		660						665					670		
Lys	Asp	Ser	Arg	Arg	Gly	Arg	Leu	Glu	Arg	Ala	Trp	Met	Asp	Gly	Ser
		675					680					685			
His	Arg	Asp	Ile	Phe	Val	Thr	Ser	Lys	Thr	Val	Leu	Trp	Pro	Asn	Gly
	690					695					700				
Leu	Ser	Leu	Asp	Ile	Pro	Ala	Gly	Arg	Leu	Tyr	Trp	Val	Asp	Ala	Phe
705					710					715					720
Tyr	Asp	Arg	Ile	Glu	Thr	Ile	Leu	Leu	Asn	Gly	Thr	Asp	Arg	Lys	Ile
			725						730					735	
Val	Tyr	Glu	Gly	Pro	Glu	Leu	Asn	His	Ala	Phe	Gly	Leu	Cys	His	His
		740						745					750		
Gly	Asn	Tyr	Leu	Phe	Trp	Thr	Glu	Tyr	Arg	Ser	Gly	Ser	Val	Tyr	Arg
		755					760					765			
Leu	Glu	Arg	Gly	Val	Ala	Gly	Ala	Pro	Pro	Thr	Val	Thr	Leu	Leu	Arg
	770					775					780				
Ser	Glu	Arg	Pro	Pro	Ile	Phe	Glu	Ile	Arg	Met	Tyr	Asp	Ala	His	Glu
785					790					795					800
Gln	Gln	Val	Gly	Thr	Asn	Lys	Cys	Arg	Val	Asn	Asn	Gly	Gly	Cys	Ser
			805						810					815	
Ser	Leu	Cys	Leu	Ala	Thr	Pro	Gly	Ser	Arg	Gln	Cys	Ala	Cys	Ala	Glu
			820					825					830		
Asp	Gln	Val	Leu	Asp	Thr	Asp	Gly	Val	Thr	Cys	Leu	Ala	Asn	Pro	Ser
		835					840					845			
Tyr	Val	Pro	Pro	Pro	Gln	Cys	Gln	Pro	Gly	Gln	Phe	Ala	Cys	Ala	Asn
	850				855						860				
Asn	Arg	Cys	Ile	Gln	Glu	Arg	Trp	Lys	Cys	Asp	Gly	Asp	Asn	Asp	Cys
865				870						875					880
Leu	Asp	Asn	Ser	Asp	Glu	Ala	Pro	Ala	Leu	Cys	His	Gln	His	Thr	Cys
			885						890					895	
Pro	Ser	Asp	Arg	Phe	Lys	Cys	Glu	Asn	Asn	Arg	Cys	Ile	Pro	Asn	Arg
			900					905					910		
Trp	Leu	Cys	Asp	Gly	Asp	Asn	Asp	Cys	Gly	Asn	Ser	Glu	Asp	Glu	Ser
		915					920					925			
Asn	Ala	Thr	Cys	Ser	Ala	Arg	Thr	Cys	Pro	Pro	Asn	Gln	Phe	Ser	Cys
	930				935						940				
Ala	Ser	Gly	Arg	Cys	Ile	Pro	Ile	Ser	Trp	Thr	Cys	Asp	Leu	Asp	Asp
945				950						955					960
Asp	Cys	Gly	Asp	Arg	Ser	Asp	Glu	Ser	Ala	Ser	Cys	Ala	Tyr	Pro	Thr
			965					970						975	

Cys Phe Pro Leu Thr Gln Phe Thr Cys Asn Asn Gly Arg Cys Ile Asn
 980 985 990
 Ile Asn Trp Arg Cys Asp Asn Asp Cys Gly Asp Asn Ser Asp
 995 1000 1005
 Glu Ala Gly Cys Ser His Ser Cys Ser Ser Thr Gln Phe Lys Cys Asn
 1010 1015 1020
 Ser Gly Arg Cys Ile Pro Glu His Trp Thr Cys Asp Gly Asp Asn Asp
 1025 1030 1035 1040
 Cys Gly Asp Tyr Ser Asp Glu Thr His Ala Asn Cys Thr Asn Gln Ala
 1045 1050 1055
 Thr Arg Pro Pro Gly Gly Cys His Ser Asp Glu Phe Gln Cys Pro Leu
 1060 1065 1070
 Asp Gly Leu Cys Ile Pro Leu Arg Trp Arg Cys Asp Gly Asp Thr Asp
 1075 1080 1085
 Cys Met Asp Ser Ser Asp Glu Lys Ser Cys Glu Gly Val Thr His Val
 1090 1095 1100
 Cys Asp Pro Asn Val Lys Phe Gly Cys Lys Asp Ser Ala Arg Cys Ile
 1105 1110 1115 1120
 Ser Lys Ala Trp Val Cys Asp Gly Asp Ser Asp Cys Glu Asp Asn Ser
 1125 1130 1135
 Asp Glu Glu Asn Cys Glu Ala Leu Ala Cys Arg Pro Pro Ser His Pro
 1140 1145 1150
 Cys Ala Asn Asn Thr Ser Val Cys Leu Pro Pro Asp Lys Leu Cys Asp
 1155 1160 1165
 Gly Lys Asp Asp Cys Gly Asp Gly Ser Asp Glu Gly Glu Leu Cys Asp
 1170 1175 1180
 Gln Cys Ser Leu Asn Asn Gly Gly Cys Ser His Asn Cys Ser Val Ala
 1185 1190 1195 1200
 Pro Gly Glu Gly Ile Val Cys Ser Cys Pro Leu Gly Met Glu Leu Gly
 1205 1210 1215
 Ser Asp Asn His Thr Cys Gln Ile Gln Ser Tyr Cys Ala Lys His Leu
 1220 1225 1230
 Lys Cys Ser Gln Lys Cys Asp Gln Asn Lys Phe Ser Val Lys Cys Ser
 1235 1240 1245
 Cys Tyr Glu Gly Trp Val Leu Glu Pro Asp Gly Glu Thr Cys Arg Ser
 1250 1255 1260
 Leu Asp Pro Phe Lys Leu Phe Ile Ile Phe Ser Asn Arg His Glu Ile
 1265 1270 1275 1280
 Arg Arg Ile Asp Leu His Lys Gly Asp Tyr Ser Val Leu Val Pro Gly
 1285 1290 1295
 Leu Arg Asn Thr Ile Ala Leu Asp Phe His Leu Ser Gln Ser Ala Leu
 1300 1305 1310
 Tyr Trp Thr Asp Ala Val Glu Asp Lys Ile Tyr Arg Gly Lys Leu Leu
 1315 1320 1325
 Asp Asn Gly Ala Leu Thr Ser Phe Glu Val Val Ile Gln Tyr Gly Leu
 1330 1335 1340
 Ala Thr Pro Glu Gly Leu Ala Val Asp Trp Ile Ala Gly Asn Ile Tyr
 1345 1350 1355 1360
 Trp Val Glu Ser Asn Leu Asp Gln Ile Glu Val Ala Lys Leu Asp Gly
 1365 1370 1375
 Thr Leu Arg Thr Thr Leu Leu Ala Gly Asp Ile Glu His Pro Arg Ala
 1380 1385 1390
 Ile Ala Leu Asp Pro Arg Asp Gly Ile Leu Phe Trp Thr Asp Trp Asp
 1395 1400 1405
 Ala Ser Leu Pro Arg Ile Glu Ala Ala Ser Met Ser Gly Ala Gly Arg
 1410 1415 1420
 Arg Thr Ile His Arg Glu Thr Gly Ser Gly Gly Cys Ala Asn Gly Leu
 1425 1430 1435 1440

Thr Val Asp Tyr Leu Glu Lys Arg Ile Leu Trp Ile Asp Ala Arg Ser
 1445 1450 1455
 Asp Ala Ile Tyr Ser Ala Arg Tyr Asp Gly Ser Gly His Met Glu Val
 1460 1465 1470
 Leu Arg Gly His Glu Phe Leu Ser His Pro Phe Ala Val Thr Leu Tyr
 1475 1480 1485
 Gly Gly Glu Val Tyr Trp Thr Asp Trp Arg Thr Asn Thr Leu Ala Lys
 1490 1495 1500
 Ala Asn Lys Trp Thr Gly His Asn Val Thr Val Val Gln Arg Thr Asn
 1505 1510 1515 1520
 Thr Gln Pro Phe Asp Leu Gln Val Tyr His Pro Ser Arg Gln Pro Met
 1525 1530 1535
 Ala Pro Asn Pro Cys Glu Ala Asn Gly Gly Arg Gly Pro Cys Ser His
 1540 1545 1550
 Leu Cys Leu Ile Asn Tyr Asn Arg Thr Val Ser Trp Ala Cys Pro His
 1555 1560 1565
 Leu Met Lys Leu His Lys Asp Asn Thr Thr Cys Tyr Glu Phe Lys Lys
 1570 1575 1580
 Phe Leu Leu Tyr Ala Arg Gln Met Glu Ile Arg Gly Val Asp Leu Asp
 1585 1590 1595 1600
 Ala Pro Tyr Tyr Asn Tyr Ile Ile Ser Phe Thr Val Pro Asp Ile Asp
 1605 1610 1615
 Asn Val Thr Val Leu Asp Tyr Asp Ala Arg Glu Gln Arg Val Tyr Trp
 1620 1625 1630
 Ser Asp Val Arg Thr Gln Ala Ile Lys Arg Ala Phe Ile Asn Gly Thr
 1635 1640 1645
 Gly Val Glu Thr Val Val Ser Ala Asp Leu Pro Asn Ala His Gly Leu
 1650 1655 1660
 Ala Val Asp Trp Val Ser Arg Asn Leu Phe Trp Thr Ser Tyr Asp Thr
 1665 1670 1675 1680
 Asn Lys Lys Gln Ile Asn Val Ala Arg Leu Asp Gly Ser Phe Lys Asn
 1685 1690 1695
 Ala Val Val Gln Gly Leu Glu Gln Pro His Gly Leu Val Val His Pro
 1700 1705 1710
 Leu Arg Gly Lys Leu Tyr Trp Thr Asp Gly Asp Asn Ile Ser Met Ala
 1715 1720 1725
 Asn Met Asp Gly Ser Asn His Thr Leu Leu Phe Ser Gly Gln Lys Gly
 1730 1735 1740
 Pro Val Gly Leu Ala Ile Asp Phe Pro Glu Ser Lys Leu Tyr Trp Ile
 1745 1750 1755 1760
 Ser Ser Gly Asn His Thr Ile Asn Arg Cys Asn Leu Asp Gly Ser Glu
 1765 1770 1775
 Leu Glu Val Ile Asp Thr Met Arg Ser Gln Leu Gly Lys Ala Thr Ala
 1780 1785 1790
 Leu Ala Ile Met Gly Asp Lys Leu Trp Trp Ala Asp Gln Val Ser Glu
 1795 1800 1805
 Lys Met Gly Thr Cys Asn Lys Ala Asp Gly Ser Gly Ser Val Val Leu
 1810 1815 1820
 Arg Asn Ser Thr Thr Leu Val Met His Met Lys Val Tyr Asp Glu Ser
 1825 1830 1835 1840
 Ile Gln Leu Glu His Glu Gly Thr Asn Pro Cys Ser Val Asn Asn Gly
 1845 1850 1855
 Asp Cys Ser Gln Leu Cys Leu Pro Thr Ser Glu Thr Thr Arg Ser Cys
 1860 1865 1870
 Met Cys Thr Ala Gly Tyr Ser Leu Arg Ser Gly Gln Gln Ala Cys Glu
 1875 1880 1885
 Gly Val Gly Ser Phe Leu Leu Tyr Ser Val His Glu Gly Ile Arg Gly
 1890 1895 1900

Ile Pro Leu Asp Pro Asn Asp Lys Ser Asp Ala Leu Val Pro Val Ser
 1905 1910 1915 1920
 Gly Thr Ser Leu Ala Val Gly Ile Asp Phe His Ala Glu Asn Asp Thr
 1925 1930 1935
 Ile Tyr Trp Val Asp Met Gly Leu Ser Thr Ile Ser Arg Ala Lys Arg
 1940 1945 1950
 Asp Gln Thr Trp Arg Glu Asp Val Val Thr Asn Gly Ile Gly Arg Val
 1955 1960 1965
 Glu Gly Ile Ala Val Asp Trp Ile Ala Gly Asn Ile Tyr Trp Thr Asp
 1970 1975 1980
 Gln Gly Phe Asp Val Ile Glu Val Ala Arg Leu Asn Gly Ser Phe Arg
 1985 1990 1995 2000
 Tyr Val Val Ile Ser Gln Gly Leu Asp Lys Pro Arg Ala Ile Thr Val
 2005 2010 2015
 His Pro Glu Lys Gly Tyr Leu Phe Trp Thr Glu Trp Gly His Tyr Pro
 2020 2025 2030
 Arg Ile Glu Arg Ser Arg Leu Asp Gly Thr Glu Arg Val Val Leu Val
 2035 2040 2045
 Asn Val Ser Ile Ser Trp Pro Asn Gly Ile Ser Val Asp Tyr Gln Gly
 2050 2055 2060
 Gly Lys Leu Tyr Trp Cys Asp Ala Arg Met Asp Lys Ile Glu Arg Ile
 2065 2070 2075 2080
 Asp Leu Glu Thr Gly Glu Asn Arg Glu Val Val Leu Ser Ser Asn Asn
 2085 2090 2095
 Met Asp Met Phe Ser Val Ser Val Phe Glu Asp Phe Ile Tyr Trp Ser
 2100 2105 2110
 Asp Arg Thr His Ala Asn Gly Ser Ile Lys Arg Gly Cys Lys Asp Asn
 2115 2120 2125
 Ala Thr Asp Ser Val Pro Leu Arg Thr Gly Ile Gly Val Gln Leu Lys
 2130 2135 2140
 Asp Ile Lys Val Phe Asn Arg Asp Arg Gln Lys Gly Thr Asn Val Cys
 2145 2150 2155 2160
 Ala Val Ala Asn Gly Gly Cys Gln Gln Leu Cys Leu Tyr Arg Gly Gly
 2165 2170 2175
 Gly Gln Arg Ala Cys Ala Cys Ala His Gly Met Leu Ala Glu Asp Gly
 2180 2185 2190
 Ala Ser Cys Arg Glu Tyr Ala Gly Tyr Leu Leu Tyr Ser Glu Arg Thr
 2195 2200 2205
 Ile Leu Lys Ser Ile His Leu Ser Asp Glu Arg Asn Leu Asn Ala Pro
 2210 2215 2220
 Val Gln Pro Phe Glu Asp Pro Glu His Met Lys Asn Val Ile Ala Leu
 2225 2230 2235 2240
 Ala Phe Asp Tyr Arg Ala Gly Thr Ser Pro Gly Thr Pro Asn Arg Ile
 2245 2250 2255
 Phe Phe Ser Asp Ile His Phe Gly Asn Ile Gln Gln Ile Asn Asp Asp
 2260 2265 2270
 Gly Ser Gly Arg Thr Thr Ile Val Glu Asn Val Gly Ser Val Glu Gly
 2275 2280 2285
 Leu Ala Tyr His Arg Gly Trp Asp Thr Leu Tyr Trp Thr Ser Tyr Thr
 2290 2295 2300
 Thr Ser Thr Ile Thr Arg His Thr Val Asp Gln Thr Arg Pro Gly Ala
 2305 2310 2315 2320
 Phe Glu Arg Glu Thr Val Ile Thr Met Ser Gly Asp Asp His Pro Arg
 2325 2330 2335
 Ala Phe Val Leu Asp Glu Cys Gln Asn Leu Met Phe Trp Thr Asn Trp
 2340 2345 2350
 Asn Glu Leu His Pro Ser Ile Met Arg Ala Ala Leu Ser Gly Ala Asn
 2355 2360 2365

Val Leu Thr Leu Ile Glu Lys Asp Ile Arg Thr Pro Asn Gly Leu Ala
2370 2375 2380
Ile Asp His Arg Ala Glu Lys Leu Tyr Phe Ser Asp Ala Thr Leu Asp
2385 2390 2395 2400
Lys Ile Glu Arg Cys Glu Tyr Asp Gly Ser His Arg Tyr Val Ile Leu
2405 2410 2415
Lys Ser Glu Pro Val His Pro Phe Gly Leu Ala Val Tyr Gly Glu His
2420 2425 2430
Ile Phe Trp Thr Asp Trp Val Arg Arg Ala Val Gln Arg Ala Asn Lys
2435 2440 2445
Tyr Val Gly Ser Asp Met Lys Leu Leu Arg Val Asp Ile Pro Gln Gln
2450 2455 2460
Pro Met Gly Ile Ile Ala Val Ala Asn Asp Thr Asn Ser Cys Glu Leu
2465 2470 2475 2480
Ser Pro Cys Arg Ile Asn Asn Gly Gly Cys Gln Asp Leu Cys Leu Leu
2485 2490 2495
Thr His Gln Gly His Val Asn Cys Ser Cys Arg Gly Gly Arg Ile Leu
2500 2505 2510
Gln Glu Asp Phe Thr Cys Arg Ala Val Asn Ser Ser Cys Arg Ala Gln
2515 2520 2525
Asp Glu Phe Glu Cys Ala Asn Gly Glu Cys Ile Ser Phe Ser Leu Thr
2530 2535 2540
Cys Asp Gly Val Ser His Cys Lys Asp Lys Ser Asp Glu Lys Pro Ser
2545 2550 2555 2560
Tyr Cys Asn Ser Arg Arg Cys Lys Lys Thr Phe Arg Gln Cys Asn Asn
2565 2570 2575
Gly Arg Cys Val Ser Asn Met Leu Trp Cys Asn Gly Val Asp Tyr Cys
2580 2585 2590
Gly Asp Gly Ser Asp Glu Ile Pro Cys Asn Lys Thr Ala Cys Gly Val
2595 2600 2605
Gly Glu Phe Arg Cys Arg Asp Gly Ser Cys Ile Gly Asn Ser Ser Arg
2610 2615 2620
Cys Asn Gln Phe Val Asp Cys Glu Asp Ala Ser Asp Glu Met Asn Cys
2625 2630 2635 2640
Ser Ala Thr Asp Cys Ser Ser Tyr Phe Arg Leu Gly Val Lys Gly Val
2645 2650 2655
Leu Phe Gln Pro Cys Glu Arg Thr Ser Leu Cys Tyr Ala Pro Ser Trp
2660 2665 2670
Val Cys Asp Gly Ala Asn Asp Cys Gly Asp Tyr Ser Asp Glu Arg Asp
2675 2680 2685
Cys Pro Gly Val Lys Arg Pro Arg Cys Pro Leu Asn Tyr Phe Ala Cys
2690 2695 2700
Pro Ser Gly Arg Cys Ile Pro Met Ser Trp Thr Cys Asp Lys Glu Asp
2705 2710 2715 2720
Asp Cys Glu Asn Gly Glu Asp Glu Thr His Cys Asn Lys Phe Cys Ser
2725 2730 2735
Glu Ala Gln Phe Glu Cys Gln Asn His Arg Cys Ile Ser Lys Gln Trp
2740 2745 2750
Leu Cys Asp Gly Ser Asp Asp Cys Gly Asp Gly Ser Asp Glu Ala Ala
2755 2760 2765
His Cys Glu Gly Lys Thr Cys Gly Pro Ser Ser Phe Ser Cys Pro Gly
2770 2775 2780
Thr His Val Cys Val Pro Glu Arg Trp Leu Cys Asp Gly Asp Lys Asp
2785 2790 2795 2800
Cys Thr Asp Gly Ala Asp Glu Ser Val Thr Ala Gly Cys Leu Tyr Asn
2805 2810 2815
Ser Thr Cys Asp Asp Arg Glu Phe Met Cys Gln Asn Arg Leu Cys Ile
2820 2825 2830

Pro Lys His Phe Val Cys Asp His Asp Arg Asp Cys Ala Asp Gly Ser
 2835 2840 2845
 Asp Glu Ser Pro Glu Cys Glu Tyr Pro Thr Cys Gly Pro Asn Glu Phe
 2850 2855 2860
 Arg Cys Ala Asn Gly Arg Cys Leu Ser Ser Arg Gln Trp Glu Cys Asp
 2865 2870 2875 2880
 Gly Glu Asn Asp Cys His Asp His Ser Asp Glu Ala Pro Lys Asn Pro
 2885 2890 2895
 His Cys Thr Ser Pro Glu His Lys Cys Asn Ala Ser Ser Gln Phe Leu
 2900 2905 2910
 Cys Ser Ser Gly Arg Cys Val Ala Glu Ala Leu Leu Cys Asn Gly Gln
 2915 2920 2925
 Asp Asp Cys Gly Asp Gly Ser Asp Glu Arg Gly Cys His Val Asn Glu
 2930 2935 2940
 Cys Leu Ser Arg Lys Leu Ser Gly Cys Ser Gln Asp Cys Glu Asp Leu
 2945 2950 2955 2960
 Lys Ile Gly Phe Lys Cys Arg Cys Arg Pro Gly Phe Arg Leu Lys Asp
 2965 2970 2975
 Asp Gly Arg Thr Cys Ala Asp Leu Asp Glu Cys Ser Thr Thr Phe Pro
 2980 2985 2990
 Cys Ser Gln Leu Cys Ile Asn Thr His Gly Ser Tyr Lys Cys Leu Cys
 2995 3000 3005
 Val Glu Gly Tyr Ala Pro Arg Gly Gly Asp Pro His Ser Cys Lys Ala
 3010 3015 3020
 Val Thr Asp Glu Glu Pro Phe Leu Ile Phe Ala Asn Arg Tyr Tyr Leu
 3025 3030 3035 3040
 Arg Lys Leu Asn Leu Asp Gly Ser Asn Tyr Thr Leu Leu Lys Gln Gly
 3045 3050 3055
 Leu Asn Asn Ala Val Ala Leu Ala Phe Asp Tyr Arg Glu Gln Met Ile
 3060 3065 3070
 Tyr Trp Thr Gly Val Thr Thr Gln Gly Ser Met Ile Arg Arg Met His
 3075 3080 3085
 Leu Asn Gly Ser Asn Val Gln Val Leu His Arg Thr Gly Leu Ser Asn
 3090 3095 3100
 Pro Asp Gly Leu Ala Val Asp Trp Val Gly Gly Asn Leu Tyr Trp Cys
 3105 3110 3115 3120
 Asp Lys Gly Arg Asp Thr Ile Glu Val Ser Lys Leu Asn Gly Ala Tyr
 3125 3130 3135
 Arg Thr Val Leu Val Ser Ser Gly Leu Arg Glu Pro Arg Ala Leu Val
 3140 3145 3150
 Val Asp Val Gln Asn Gly Tyr Leu Tyr Trp Thr Asp Trp Gly Asp His
 3155 3160 3165
 Ser Leu Ile Gly Arg Ile Gly Met Asp Gly Ser Gly Arg Ser Ile Ile
 3170 3175 3180
 Val Asp Thr Lys Ile Thr Trp Pro Asn Gly Leu Thr Val Asp Tyr Val
 3185 3190 3195 3200
 Thr Glu Arg Ile Tyr Trp Ala Asp Ala Arg Glu Asp Tyr Ile Glu Phe
 3205 3210 3215
 Ala Ser Leu Asp Gly Ser Asn Arg His Val Val Leu Ser Gln Asp Ile
 3220 3225 3230
 Pro His Ile Phe Ala Leu Thr Leu Phe Glu Asp Tyr Val Tyr Trp Thr
 3235 3240 3245
 Asp Trp Glu Thr Lys Ser Ile Asn Arg Ala His Lys Thr Thr Gly Ala
 3250 3255 3260
 Asn Lys Thr Leu Leu Ile Ser Thr Leu His Arg Pro Met Asp Leu His
 3265 3270 3275 3280
 Val Phe His Ala Leu Arg Gln Pro Asp Val Pro Asn His Pro Cys Lys
 3285 3290 3295

Val Asn Asn Gly Gly Cys Ser Asn Leu Cys Leu Leu Ser Pro Gly Gly
 3300 3305 3310
 Gly His Lys Cys Ala Cys Pro Thr Asn Phe Tyr Leu Gly Gly Asp Gly
 3315 3320 3325
 Arg Thr Cys Val Ser Asn Cys Thr Ala Ser Gln Phe Val Cys Lys Asn
 3330 3335 3340
 Asp Lys Cys Ile Pro Phe Trp Trp Lys Cys Asp Thr Glu Asp Asp Cys
 3345 3350 3355 3360
 Gly Asp His Ser Asp Glu Pro Pro Asp Cys Pro Glu Phe Lys Cys Arg
 3365 3370 3375
 Pro Gly Gln Phe Gln Cys Ser Thr Gly Ile Cys Thr Asn Pro Ala Phe
 3380 3385 3390
 Ile Cys Asp Gly Asp Asn Asp Cys Gln Asp Asn Ser Asp Glu Ala Asn
 3395 3400 3405
 Cys Asp Ile His Val Cys Leu Pro Ser Gln Phe Lys Cys Thr Asn Thr
 3410 3415 3420
 Asn Arg Cys Ile Pro Gly Ile Phe Arg Cys Asn Gly Gln Asp Asn Cys
 3425 3430 3435 3440
 Gly Asp Gly Glu Asp Glu Arg Asp Cys Pro Glu Val Thr Cys Ala Pro
 3445 3450 3455
 Asn Gln Phe Gln Cys Ser Ile Thr Lys Arg Cys Ile Pro Arg Val Trp
 3460 3465 3470
 Val Cys Asp Arg Asp Asn His Cys Val Asp Gly Ser Asp Glu Pro Ala
 3475 3480 3485
 Asn Cys Thr Gln Met Thr Cys Gly Val Asp Glu Phe Arg Cys Lys Asp
 3490 3495 3500
 Ser Gly Arg Cys Ile Pro Ala Arg Trp Lys Cys Asp Gly Glu Asp Asp
 3505 3510 3515 3520
 Cys Gly Asp Gly Ser Asp Glu Pro Lys Glu Glu Cys Asp Glu Arg Thr
 3525 3530 3535
 Cys Glu Pro Tyr Gln Phe Arg Cys Lys Asn Asn Arg Cys Val Pro Gly
 3540 3545 3550
 Arg Trp Gln Cys Asp Tyr Asp Asn Asp Cys Gly Asp Asn Ser Asp Glu
 3555 3560 3565
 Glu Ser Cys Thr Pro Arg Pro Cys Ser Glu Ser Glu Phe Phe Cys Ala
 3570 3575 3580
 Asn Gly Arg Cys Ile Ala Gly Arg Trp Lys Cys Asp Gly Asp His Asp
 3585 3590 3595 3600
 Cys Ala Asp Gly Ser Asp Glu Lys Asp Cys Thr Pro Arg Cys Asp Met
 3605 3610 3615
 Asp Gln Phe Gln Cys Lys Ser Gly His Cys Ile Pro Leu Arg Trp Pro
 3620 3625 3630
 Cys Asp Ala Asp Ala Asp Cys Met Asp Gly Ser Asp Glu Glu Ala Cys
 3635 3640 3645
 Gly Thr Gly Val Arg Thr Cys Pro Leu Asp Glu Phe Gln Cys Asn Asn
 3650 3655 3660
 Thr Leu Cys Lys Pro Leu Ala Trp Lys Cys Asp Gly Glu Asp Asp Cys
 3665 3670 3675 3680
 Gly Asp Asn Ser Asp Glu Asn Pro Glu Glu Cys Ala Arg Phe Ile Cys
 3685 3690 3695
 Pro Pro Asn Arg Pro Phe Arg Cys Lys Asn Asp Arg Val Cys Leu Trp
 3700 3705 3710
 Ile Gly Arg Gln Cys Asp Gly Val Asp Asn Cys Gly Asp Gly Thr Asp
 3715 3720 3725
 Glu Glu Asp Cys Glu Pro Pro Thr Ala Gln Asn Pro His Cys Lys Asp
 3730 3735 3740
 Lys Lys Glu Phe Leu Cys Arg Asn Gln Arg Cys Leu Ser Ser Ser Leu
 3745 3750 3755 3760

Arg Cys Asn Met Phe Asp Asp Cys Gly Asp Gly Ser Asp Glu Glu Asp
 3765 3770 3775
 Cys Ser Ile Asp Pro Lys Leu Thr Ser Cys Ala Thr Asn Ala Ser Met
 3780 3785 3790
 Cys Gly Asp Glu Ala Arg Cys Val Arg Thr Glu Lys Ala Ala Tyr Cys
 3795 3800 3805
 Ala Cys Arg Ser Gly Phe His Thr Val Pro Gly Gln Pro Gly Cys Gln
 3810 3815 3820
 Asp Ile Asn Glu Cys Leu Arg Phe Gly Thr Cys Ser Gln Leu Trp Asn
 3825 3830 3835 3840
 Lys Pro Lys Gly Gly His Leu Cys Ser Cys Ala Arg Asn Phe Met Lys
 3845 3850 3855
 Thr His Asn Thr Cys Lys Ala Glu Gly Ser Glu Tyr Gln Val Leu Tyr
 3860 3865 3870
 Ile Ala Asp Asp Asn Glu Ile Arg Ser Leu Phe Pro Gly His Pro His
 3875 3880 3885
 Ser Ala Tyr Glu Gln Thr Phe Gln Gly Asp Glu Ser Val Arg Ile Asp
 3890 3895 3900
 Ala Met Asp Val His Val Lys Ala Gly Arg Val Tyr Trp Thr Asn Trp
 3905 3910 3915 3920
 His Thr Gly Thr Ile Ser Tyr Arg Ser Leu Pro Pro Ala Ala Pro Pro
 3925 3930 3935
 Thr Thr Ser Asn Arg His Arg Arg Gln Ile Asp Arg Gly Val Thr His
 3940 3945 3950
 Leu Asn Ile Ser Gly Leu Lys Met Pro Arg Gly Ile Ala Ile Asp Trp
 3955 3960 3965
 Val Ala Gly Asn Val Tyr Trp Thr Asp Ser Gly Arg Asp Val Ile Glu
 3970 3975 3980
 Val Ala Gln Met Lys Gly Glu Asn Arg Lys Thr Leu Ile Ser Gly Met
 3985 3990 3995 4000
 Ile Asp Glu Pro His Ala Ile Val Val Asp Pro Leu Arg Gly Thr Met
 4005 4010 4015
 Tyr Trp Ser Asp Trp Gly Asn His Pro Lys Ile Glu Thr Ala Ala Met
 4020 4025 4030
 Asp Gly Thr Leu Arg Glu Thr Leu Val Gln Asp Asn Ile Gln Trp Pro
 4035 4040 4045
 Thr Gly Leu Ala Val Asp Tyr His Asn Glu Arg Leu Tyr Trp Ala Asp
 4050 4055 4060
 Ala Lys Leu Ser Val Ile Gly Ser Ile Arg Leu Asn Gly Thr Asp Pro
 4065 4070 4075 4080
 Ile Val Ala Ala Asp Ser Lys Arg Gly Leu Ser His Pro Phe Ser Ile
 4085 4090 4095
 Asp Val Phe Glu Asp Tyr Ile Tyr Gly Val Thr Tyr Ile Asn Asn Arg
 4100 4105 4110
 Val Phe Lys Ile His Lys Phe Gly His Ser Pro Leu Tyr Asn Leu Thr
 4115 4120 4125
 Gly Gly Leu Ser His Ala Ser Asp Val Val Leu Tyr His Gln His Lys
 4130 4135 4140
 Gln Pro Glu Val Thr Asn Pro Cys Asp Arg Lys Lys Cys Glu Trp Leu
 4145 4150 4155 4160
 Cys Leu Leu Ser Pro Ser Gly Pro Val Cys Thr Cys Pro Asn Gly Lys
 4165 4170 4175
 Arg Leu Asp Asn Gly Thr Cys Val Pro Val Pro Ser Pro Thr Pro Pro
 4180 4185 4190
 Pro Asp Ala Pro Arg Pro Gly Thr Cys Thr Leu Gln Cys Phe Asn Gly
 4195 4200 4205
 Gly Ser Cys Phe Leu Asn Ala Arg Arg Gln Pro Lys Cys Arg Cys Gln
 4210 4215 4220

Pro Arg Tyr Thr Gly Asp Lys Cys Glu Leu Asp Gln Cys Trp Glu Tyr
 4225 4230 4235 4240
 Cys His Asn Gly Gly Thr Cys Ala Ala Ser Pro Ser Gly Met Pro Thr
 4245 4250 4255
 Cys Arg Cys Pro Thr Gly Phe Thr Gly Pro Lys Cys Thr Ala Gln Val
 4260 4265 4270
 Cys Ala Gly Tyr Cys Ser Asn Asn Ser Thr Cys Thr Val Asn Gln Gly
 4275 4280 4285
 Asn Gln Pro Gln Cys Arg Cys Leu Pro Gly Phe Leu Gly Asp Arg Cys
 4290 4295 4300
 Gln Tyr Arg Gln Cys Ser Gly Phe Cys Glu Asn Phe Gly Thr Cys Gln
 4305 4310 4315 4320
 Met Ala Ala Asp Gly Ser Arg Gln Cys Arg Cys Thr Val Tyr Phe Glu
 4325 4330 4335
 Gly Pro Arg Cys Glu Val Asn Lys Cys Ser Arg Cys Leu Gln Gly Ala
 4340 4345 4350
 Cys Val Val Asn Lys Gln Thr Gly Asp Val Thr Cys Asn Cys Thr Asp
 4355 4360 4365
 Gly Arg Val Ala Pro Ser Cys Leu Thr Cys Ile Asp His Cys Ser Asn
 4370 4375 4380
 Gly Gly Ser Cys Thr Met Asn Ser Lys Met Met Pro Glu Cys Gln Cys
 4385 4390 4395 4400
 Pro Pro His Met Thr Gly Pro Arg Cys Gln Glu Gln Val Val Ser Gln
 4405 4410 4415
 Gln Gln Pro Gly His Met Ala Ser Ile Leu Ile Pro Leu Leu Leu Leu
 4420 4425 4430
 Leu Leu Leu Leu Leu Val Ala Gly Val Val Phe Trp Tyr Lys Arg Arg
 4435 4440 4445
 Val Arg Gly Ala Lys Gly Phe Gln His Gln Arg Met Thr Asn Gly Ala
 4450 4455 4460
 Met Asn Val Glu Ile Gly Asn Pro Thr Tyr Lys Met Tyr Glu Gly Gly
 4465 4470 4475 4480
 Glu Pro Asp Asp Val Gly Gly Leu Leu Asp Ala Asp Phe Ala Leu Asp
 4485 4490 4495
 Pro Asp Lys Pro Thr Asn Phe Thr Asn Pro Val Tyr Ala Thr Leu Tyr
 4500 4505 4510
 Met Gly Gly His Gly Ser Arg His Ser Leu Ala Ser Thr Asp Glu Lys
 4515 4520 4525
 Arg Glu Leu Leu Gly Arg Gly Pro Glu Asp Glu Ile Gly Asp Pro Leu
 4530 4535 4540
 Ala
 4545

<210> 3
 <211> 4577
 <212> DNA
 <213> Homo sapiens

<400> 3
 gctacaatcc atctggtctc ctccagctcc ttctttctgc aacatgggga agaacaaact 60
 ccttcattcca agtctggttc ttctcctctt ggtcctcctg cccacagacg cctcagtctc 120
 tggaaaaccg cagtatatgg ttctggtccc ctccctgctc cacactgaga ccactgagaa 180
 gggctgtgtc cttctgagct acctgaatga gacagtgact gtaagtgtt ccttggagtc 240
 tgtcagggga aacaggagcc tcttctactga cctggaggcg gagaatgacg tactccactg 300
 tgtcgccttc gctgtcccaa agtcttcatc caatgaggag gtaatgttcc tctactgtcca 360
 agtgaaagga ccaacccaag aatttaagaa gcggaccaca gtgatgggta agaacgagga 420
 cagtctggtc tttgtccaga cagacaaatc aatctacaaa ccagggcaga cagtgaatt 480
 tcgtgttgc tccatggatg aaaactttca cccctgaat gagttgattc cactagtata 540

cattcaggat	cccaaaggaa	atcgcacgc	acaatggcag	agtttccagt	tagaggggtg	600
cctcaagcaa	ttttcttttc	ccctctcatc	agagcccttc	caggggtcct	acaaggtggt	660
ggtacagaag	aaatcagggtg	gaaggacaga	gcaccctttc	accgtggagg	aatttgttct	720
tcccaagttt	gaagtacaag	taacagtgcc	aaagataatc	accatcttgg	aagaagagat	780
gaatgtatca	gtgtgtggcc	tatacacata	tgggaagcct	gtccctggac	atgtgactgt	840
gagcatttgc	agaaagtata	gtgacgcttc	cgactgccac	ggtgaagatt	cacaggcttt	900
ctgtgagaaa	ttcagtggac	agctaaacag	ccatggctgc	ttctatcagc	aagtaaaaaac	960
caaggtcttc	cagctgaaga	ggaaggagta	tgaatgaaa	cttcacactg	aggcccagat	1020
ccaagaagaa	ggaacagtgg	tggaaattgac	tgggaaggcag	tccagtgaaa	tcacaagaac	1080
cataaccaaa	ctctcatttg	tgaaagtggg	ctcacacttt	cgacagggaa	ttcccttctt	1140
tgggcagggtg	cgcctagtag	atgggaaagg	cgtccctata	ccaaataaag	tcataattcat	1200
cagaggaaat	gaagcaaat	attactccaa	tgtaccacg	gatgagcatg	gccttgtaga	1260
gttctctatc	aacaccacca	acgttatggg	tacctctctt	actgttaggg	tcaattacaa	1320
ggatcgtagt	ccctgttacg	gctaccagtg	ggtgtcagaa	gaacacgaag	aggcacatca	1380
cactgcttat	cttgtgttct	ccccaaagcaa	gagctttgtc	caccttgagc	ccatgtctca	1440
tgaactaccc	tgtggccata	ctcagacagt	ccaggcacat	tatatcttga	atggaggcac	1500
cctgctgggg	ctgaagaagc	tctcctttta	ttatctgata	atggcaaagg	gaggcattgt	1560
ccgaactggg	actcatggac	tgcttgtgaa	gcaggaagac	atgaagggcc	atttttccat	1620
ctcaatccct	gtgaagtcag	acattgctcc	tgtcgctcgg	ttgctcatct	atgctgtttt	1680
acctaccggg	gacgtgattg	gggattctgc	aaaatatgat	gttgaaaatt	gtctggccaa	1740
caaggtggat	ttgagcttca	gcccatacaca	aagtctccca	gcctcacacg	cccacctgcg	1800
agtcacagcg	gctcctcagt	ccgtctgcgc	cctccgtgct	gtggaccaaa	gcgtgctgct	1860
catgaagcct	gatgctgagc	tctcggcgtc	ctcggtttac	aacctgctac	cagaaaagga	1920
cctcactggc	ttccctgggc	ctttgaatga	ccaggacgat	gaagactgca	tcaatcgtca	1980
taatgtctat	attaatggaa	tcacatatac	tccagtatca	agtacaaatg	aaaaggatat	2040
gtacagcttc	ctagaggaca	tgggcttaaa	ggcattcacc	aactcaaaga	ttcgtaaacc	2100
caaaatgtgt	ccacagcttc	aacagtatga	aatgcatgga	cctgaaggctc	tacgtgtagg	2160
tttttatgag	tcagatgtaa	tgggaagagg	ccatgcacgc	ctgggtgcatg	ttgaagagcc	2220
tcacacggag	accgtacgaa	agtacttccc	tgagacatgg	atctgggatt	tgggtggtgt	2280
aaactcagca	ggggtggctg	aggtaggagt	aacagtcctc	gacaccatca	ccgagtggaa	2340
ggcagggggc	ttctgcctgt	ctgaagatgc	tggacttggt	atctcttcca	ctgcctctct	2400
ccgagccttc	cagcccttct	ttgtggagct	tacaatgcct	tactctgtga	ttcgtggaga	2460
ggccttcaca	ctcaaggcca	cggtcctaaa	ctaccttccc	aaatgcatcc	gggtcagtgt	2520
gcagctggaa	gcctctcccg	ccttccttgc	tgtcccagtg	gagaaggaaac	aagcgcttca	2580
ctgcatctgt	gcaaacgggc	ggcaaactgt	gtcctgggca	gtaaccccaa	agtcattagg	2640
aaatgtgaat	ttcactgtga	gcgcagaggc	actagagtct	caagagctgt	gtgggactga	2700
ggtgccttca	gttccctgaac	acggaaggaa	agacacagtc	atcaagcctc	tgttggttga	2760
acctgaagga	ctagagaagg	aaacaacatt	caactcccta	ctttgtccat	caggtggtga	2820
ggtttctgaa	gaattatccc	tgaactgccc	accaaatgtg	gtagaagaat	ctgcccgagc	2880
ttctgtctca	gttttgggag	acatattagg	ctctgccatg	caaaacacac	aaaatcttct	2940
ccagatgccc	tatggctgtg	gagagcagaa	tatggtcctc	tttgcctcta	acatctatgt	3000
actggattat	ctaaatgaaa	cacagcagct	tactccagag	gtcaagtcca	aggccattgg	3060
ctatctcaac	actggttacc	agagacagtt	gaactacaaa	cactatgatg	gctcctacag	3120
cacctttggg	gagcgatatg	gcaggaacca	gggcaacacc	tggctcacag	cctttgttct	3180
gaagactttt	gccccagctc	gagcctacat	cttcacgat	gaagcacaca	ttacccaagc	3240
cctcatatgg	ctctcccaga	ggcagaagga	caatggctgt	ttcaggagct	ctgggtcact	3300
gctcaacaat	gccataaagg	gaggagtaga	agatgaagtg	accctctccg	cctatatcac	3360
catgcgccct	ctggagattc	ctctcacagt	cactcaccct	gttgtccgca	atgccctgtt	3420
ttgcctggag	tcagcctgga	agacagcaca	agaaggggac	catggcagcc	atgtatatac	3480
caaagcactg	ctggcctatg	cttttgccct	ggcaggtaac	caggacaaga	ggaagggaagt	3540
actcaagtca	cttaatgagg	aagctgtgaa	gaaagacaac	tctgtccatt	gggagcgccc	3600
tcagaaaccc	aaggcaccag	tggggcattt	ttacgaaccc	caggctccct	ctgctgaggt	3660
ggagatgaca	tcctatgtgc	tcctcgctta	tctcacggcc	cagccagccc	caacctcgga	3720
ggacctgacc	tctgcaacca	acatcgtgaa	gtggatcacg	aagcagcaga	atgcccaggg	3780
cggtttctcc	tccaccaggg	acacagtggt	ggctctccat	gctctgtcca	aatatggagc	3840
cgccacattt	accaggactg	ggaaggctgc	acaggtgact	atccagtctt	cagggacatt	3900
ttccagcaaa	ttccaagtgg	acaacaacaa	tcgcctgtta	ctgcagcagg	tctcattgcc	3960
agagctgcct	ggggaatata	gcattgaaagt	gacaggagaa	ggatgtgtct	acctccagac	4020

ctccttgaaa	tacaatat	tcccagaaaa	ggaagagttc	ccctttgctt	taggagtgca	4080
gactctgcct	caaacttg	atgaacccaa	agcccacacc	agcttccaaa	tctccctaag	4140
tgtcagttac	acagggagcc	gctctgcctc	caacatggcg	atcggtgatg	tgaagatggg	4200
ctctggcttc	attccctga	agccaacagt	gaaaatgctt	gaaagatcta	accatgtgag	4260
ccggacagaa	gtcagcagca	accatgtctt	gatttacctt	gataaggtgt	caaatacagac	4320
actgagcttg	ttcttcacgg	ttctgcaaga	tgtcccagta	agagatctca	aaccagccat	4380
agtgaaagtc	tatgattact	acgagacgga	tgagtttgca	atcgctgagt	acaatgctcc	4440
ttgcagcaaa	gatcttggaa	atgcttgaag	accacaaggc	tgaaaagtgc	tttgctggag	4500
tcctgttctc	tgagctccac	agaagacacg	tgtttttgta	tctttaaaga	cttgatgaat	4560
aaacactttt	tctggtc					4577

<210> 4

<211> 4422

<212> DNA

<213> Homo sapiens

<400> 4

atggggaaga	acaaactcct	tcatccaagt	ctggttcttc	tcctcttggt	cctcctgccc	60
acagacgcct	cagtctctgg	aaaaccgcag	tatatggttc	tggtcccctc	cctgctccac	120
actgagacca	ctgagaaggg	ctgtgtcctt	ctgagctacc	tgaatgagac	agtgactgta	180
agtgtctcct	tggagtctgt	caggggaaac	aggagcctct	tactgacct	ggaggcggag	240
aatgacgtac	tccactgtgt	cgccttcgct	gtcccaaagt	cttcatccaa	tgaggaggta	300
atgttccctca	ctgtccaagt	gaaaggacca	acccaagaat	ttaagaagcg	gaccacagt	360
atggttaaga	acgaggacag	tctggtcttt	gtccagacag	acaaatcaat	ctacaaacca	420
gggcagacag	tgaaatttcg	tgttgtctcc	atggatgaaa	actttcaccc	cctgaatgag	480
ttgattccac	tagtatacat	tcaggatccc	aaaggaaatc	gcatacgaca	atggcagagt	540
ttccagttag	aggggtggcct	caagcaattt	tcttttcccc	tctcatcaga	gcccttccag	600
ggctcctaca	aggtggtggt	acagaagaaa	tcaggtggaa	ggacagagca	ccctttcacc	660
gtggaggaat	ttgttcttcc	caagtttgaa	gtacaagtaa	cagtgccaaa	gataatcacc	720
atcttggaag	aagagatgaa	tgtatcagtg	tgtggcctat	acacatatgg	gaagcctgtc	780
cctggacatg	tgactgtgag	catttgacaga	aagtatagtg	acgcttcgga	ctgccacggg	840
gaagattcac	aggttttctg	tgagaaattc	agtggacagc	taaacagcca	tggtgtcttc	900
tatcagcaag	taaaaaccaa	ggtcttccag	ctgaagagga	aggagtatga	aatgaaactt	960
cacactgagg	cccagatcca	agaagaagga	acagtgggtg	aattgactgg	aaggcagtcc	1020
agtgaatca	caagaaccat	aaccaaactc	tcatttgtga	aagtggactc	acactttcga	1080
caggggaattc	ccttctttgg	gcagggtgcg	ctagtagatg	ggaaaggcgt	ccctatacca	1140
aataaagtca	tattcatcag	aggaaatgaa	gcaaactatt	actccaatgc	taccacggat	1200
gagcatggcc	ttgtacagtt	ctctatcaac	accaccaacg	ttatgggtac	ctctcttact	1260
gttagggcca	attacaagga	tcgtagtccc	tgttacggct	accagtgggt	gtcagaagaa	1320
cacgaagagg	cacatcacac	tgttatctct	gtgttctccc	caagcaagag	ccttgtccac	1380
cttgagccca	tgtctcatga	actaccctgt	ggccatactc	agacagtcca	ggcacattat	1440
attctgaatg	gaggcaccct	gctggggctg	aagaagctct	ccttttatta	tctgataatg	1500
gcaaagggag	gcattgtccg	aactggggact	catggactgc	ttgtgaagca	ggaagacatg	1560
aagggccatt	tttccatctc	aatccctgtg	aagtcagaca	ttgtctctgt	cgctcgggtg	1620
ctcatctatg	ctgttttacc	taccggggac	gtgattgggg	attctgcaaa	atatgatgtt	1680
gaaaattgtc	tggccaacaa	ggtggatttg	agcttcagcc	catcacaag	tctcccagcc	1740
tcacacgccc	acctgcgagt	cacagcggct	cctcagtcct	tctgcgccct	ccgtgctgtg	1800
gaccaagcgg	tgctgtcat	gaagcctgat	gctgagctct	cggcgctcct	ggtttacaac	1860
ctgctaccag	aaaaggacct	cactggcttc	cctgggcctt	tgaatgacca	ggacgatgaa	1920
gactgcatac	atcgtcataa	tgtctatatt	aatggaatca	catatactcc	agtatcaagt	1980
acaaatgaaa	aggatatgta	cagcttccta	gaggacatgg	gcttaaaggc	attcaccaac	2040
tcaaagattc	gtaaacccaa	aatgtgtcca	cagcttcaac	agtatgaaat	gcattggacct	2100
gaaggtctac	gtgtagggtt	ttatgagtca	gatgtaatgg	gaagaggcca	tgcacgcctg	2160
gtgcatgttg	aagagcctca	cacggagacc	gtacgaaagt	acttccttga	gacatggatc	2220
tgggatttgg	tgggtgtaaa	ctcagcaggg	gtggctgagg	taggagtaac	agtccctgac	2280
accatcaccg	agtggaaagg	aggggccttc	tgcctgtctg	aagatgctgg	acttggtatc	2340
tcttccactg	cctctctccg	agccttccag	cccttctttg	tggagcttac	aatgccttac	2400
tctgtgattc	gtggagaggc	cttcacactc	aaggccacgg	tcctaaacta	ccttcccaaa	2460

tgc	atc	ccg	gg	tcag	tgt	gca	gct	gga	agcc	tct	cccg	cct	tcct	tgt	gt	ccc	agt	ggag	2520		
aag	gaaca	aag	cgc	ctc	act	g	cat	ctg	tgt	gca	aac	ggg	cgg	aa	act	gt	gtc	ctg	ggc	agta	2580
accc	caaa	agt	catt	agg	aaa	tgt	ga	att	tc	act	gt	gag	c	gag	agg	ca	ct	ca	aa	2640	
gag	ctgt	gtg	gg	act	gag	gt	gc	ttc	ag	tt	cct	ga	ac	g	ga	agg	aa	ga	ca	agtc	2700
aag	cct	ctgt	tg	gt	ga	acc	tga	agg	acta	gaga	agg	aaa	ca	ac	att	caa	ct	cc	ta	ctt	2760
tgt	ccat	cag	gt	gt	gag	gt	tt	ct	ga	aa	tt	at	cc	ct	ga	aa	act	g	cc	acc	2820
ga	aga	atct	ccc	gag	cttc	tgt	ct	cag	tt	tt	ggg	ga	ga	ca	tatt	agg	ctc	tg	cc	atg	2880
aa	ca	ca	aaa	at	ctt	ct	cca	gat	gcc	ctat	gg	ct	gt	gg	ag	caga	aat	at	gg	tc	2940
g	ct	c	ta	aca	t	ct	at	gt	act	gg	att	at	ct	a	at	gaa	ac	ac	g	ct	3000
aa	gt	cca	agg	cc	att	gg	cta	t	ct	ca	ac	act	gg	tt	acc	aga	g	ac	ag	tt	3060
ta	tg	at	gg	ct	c	ta	cag	ca	ctt	tg	gg	gg	gag	cg	at	at	gg	ca	gg	a	3120
ct	ca	cag	cct	tt	gt	ct	gaa	g	act	ttt	g	cc	ca	ag	ct	gag	c	ct	ac	at	3180
gc	ac	ac	atta	cc	ca	ag	cc	ct	ca	ta	t	gg	ct	c	ca	gag	gc	aga	agg	acaa	3240
agg	ag	ct	ctg	gg	ta	c	act	g	ct	ca	aa	at	g	cc	ata	aa	gg	gag	ga	ta	3300
ct	ct	cc	gc	ct	at	at	ca	cc	at	cg	cc	ct	ct	g	gag	at	ct	ct	c	ac	3360
gt	cc	g	ca	at	g	ct	gt	tt	tg	c	ct	gg	ag	t	ca	g	ct	ca	g	ca	3420
gg	cag	cc	atg	ta	ta	ta	cc	aa	ag	ca	ct	g	ct	g	ct	at	g	ct	tt	tg	3480
ga	ca	ag	ag	ga	ag	ta	ct	ca	ag	ta	ct	ct	aa	tg	ag	ga	ag	ct	gt	ga	3540
gt	cc	att	gg	ag	cg	cc	ct	ca	ga	ac	cc	ca	ag	gc	ac	ca	gt	gg	gg	catt	3600
g	ct	cc	ct	ctg	ct	gag	gg	t	ga	g	ga	g	ga	g	at	g	ac	at	cc	ct	3660
cc	ag	cc	caa	cc	tc	gg	ag	ga	g	at	g	ac	ct	ct	g	ca	ac	ca	ca	g	3720
cag	ca	aat	g	cc	ag	gg	c	gg	g	tt	ct	ct	cc	cc	ag	ga	ca	ag	tt	gg	3780
ct	gt	cc	aa	at	at	gg	ag	cc	gc	gc	ca	ct	ta	cc	ag	gg	ct	gc	ac	ac	3840
ca	gt	ct	tc	ag	gg	ac	att	tt	tc	ca	g	caa	at	tc	ca	ag	ta	gg	ac	ac	3900
cag	ca	gg	tt	ct	catt	gg	cc	aga	g	ct	g	cc	tg	gg	ga	ata	cag	ca	g	ac	3960
tgt	gt	ct	ta	cc	tc	ag	ac	ct	tc	ct	tg	aa	at	ac	aa	ta	tt	ct	cc	ca	4020
ttt	g	ct	tt	ag	gag	tc	ag	ac	tc	ct	g	cc	ta	aa	act	gt	gat	g	ac	cc	4080
tt	cc	aa	at	ct	cc	ta	ag	tg	g	ca	gt	ta	ca	ca	gg	g	cc	gc	ct	ca	4140
gt	t	gat	gt	ga	ag	at	gg	gt	ct	tc	tg	gt	tc	att	ccc	ct	ga	ag	gc	ca	4200
ag	at	ct	ta	acc	at	gt	gag	ccg	g	ac	ga	ag	tc	ag	ca	ga	acc	at	gt	ct	4260
aag	gt	gt	caa	at	ca	g	ac	act	gag	ct	tt	gt	tt	tc	ac	gg	tt	tc	tg	ca	4320
gat	ct	caa	ac	ca	gc	ca	ta	agt	gaa	ag	tc	ta	g	att	act	ac	g	ac	gg	at	4380
g	ct	gag	ta	ca	at	gt	ct	ct	tg	ca	aa	ag	at	ct	tg	gaa	aat	g	ct		4422

<210> 5
 <211> 1474
 <212> PRT
 <213> Homo sapiens

<400> 5
 Met Gly Lys Asn Lys Leu Leu His Pro Ser Leu Val Leu Leu Leu Leu
 1 5 10 15
 Val Leu Leu Pro Thr Asp Ala Ser Val Ser Gly Lys Pro Gln Tyr Met
 20 25 30
 Val Leu Val Pro Ser Leu Leu His Thr Glu Thr Thr Glu Lys Gly Cys
 35 40 45
 Val Leu Leu Ser Tyr Leu Asn Glu Thr Val Thr Val Ser Ala Ser Leu
 50 55 60
 Glu Ser Val Arg Gly Asn Arg Ser Leu Phe Thr Asp Leu Glu Ala Glu
 65 70 75 80
 Asn Asp Val Leu His Cys Val Ala Phe Ala Val Pro Lys Ser Ser Ser
 85 90 95
 Asn Glu Glu Val Met Phe Leu Thr Val Gln Val Lys Gly Pro Thr Gln
 100 105 110
 Glu Phe Lys Lys Arg Thr Thr Val Met Val Lys Asn Glu Asp Ser Leu
 115 120 125
 Val Phe Val Gln Thr Asp Lys Ser Ile Tyr Lys Pro Gly Gln Thr Val
 130 135 140

Lys	Phe	Arg	Val	Val	Ser	Met	Asp	Glu	Asn	Phe	His	Pro	Leu	Asn	Glu	145	150	155	160
Leu	Ile	Pro	Leu	Val	Tyr	Ile	Gln	Asp	Pro	Lys	Gly	Asn	Arg	Ile	Ala	165	170	175	
Gln	Trp	Gln	Ser	Phe	Gln	Leu	Glu	Gly	Gly	Leu	Lys	Gln	Phe	Ser	Phe	180	185	190	
Pro	Leu	Ser	Ser	Glu	Pro	Phe	Gln	Gly	Ser	Tyr	Lys	Val	Val	Val	Gln	195	200	205	
Lys	Lys	Ser	Gly	Gly	Arg	Thr	Glu	His	Pro	Phe	Thr	Val	Glu	Glu	Phe	210	215	220	
Val	Leu	Pro	Lys	Phe	Glu	Val	Gln	Val	Thr	Val	Pro	Lys	Ile	Ile	Thr	225	230	235	240
Ile	Leu	Glu	Glu	Glu	Met	Asn	Val	Ser	Val	Cys	Gly	Leu	Tyr	Thr	Tyr	245	250	255	
Gly	Lys	Pro	Val	Pro	Gly	His	Val	Thr	Val	Ser	Ile	Cys	Arg	Lys	Tyr	260	265	270	
Ser	Asp	Ala	Ser	Asp	Cys	His	Gly	Glu	Asp	Ser	Gln	Ala	Phe	Cys	Glu	275	280	285	
Lys	Phe	Ser	Gly	Gln	Leu	Asn	Ser	His	Gly	Cys	Phe	Tyr	Gln	Gln	Val	290	295	300	
Lys	Thr	Lys	Val	Phe	Gln	Leu	Lys	Arg	Lys	Glu	Tyr	Glu	Met	Lys	Leu	305	310	315	320
His	Thr	Glu	Ala	Gln	Ile	Gln	Glu	Glu	Gly	Thr	Val	Val	Glu	Leu	Thr	325	330	335	
Gly	Arg	Gln	Ser	Ser	Glu	Ile	Thr	Arg	Thr	Ile	Thr	Lys	Leu	Ser	Phe	340	345	350	
Val	Lys	Val	Asp	Ser	His	Phe	Arg	Gln	Gly	Ile	Pro	Phe	Phe	Gly	Gln	355	360	365	
Val	Arg	Leu	Val	Asp	Gly	Lys	Gly	Val	Pro	Ile	Pro	Asn	Lys	Val	Ile	370	375	380	
Phe	Ile	Arg	Gly	Asn	Glu	Ala	Asn	Tyr	Tyr	Ser	Asn	Ala	Thr	Thr	Asp	385	390	395	400
Glu	His	Gly	Leu	Val	Gln	Phe	Ser	Ile	Asn	Thr	Thr	Asn	Val	Met	Gly	405	410	415	
Thr	Ser	Leu	Thr	Val	Arg	Val	Asn	Tyr	Lys	Asp	Arg	Ser	Pro	Cys	Tyr	420	425	430	
Gly	Tyr	Gln	Trp	Val	Ser	Glu	Glu	His	Glu	Glu	Ala	His	His	Thr	Ala	435	440	445	
Tyr	Leu	Val	Phe	Ser	Pro	Ser	Lys	Ser	Phe	Val	His	Leu	Glu	Pro	Met	450	455	460	
Ser	His	Glu	Leu	Pro	Cys	Gly	His	Thr	Gln	Thr	Val	Gln	Ala	His	Tyr	465	470	475	480
Ile	Leu	Asn	Gly	Gly	Thr	Leu	Leu	Gly	Leu	Lys	Lys	Leu	Ser	Phe	Tyr	485	490	495	
Tyr	Leu	Ile	Met	Ala	Lys	Gly	Gly	Ile	Val	Arg	Thr	Gly	Thr	His	Gly	500	505	510	
Leu	Leu	Val	Lys	Gln	Glu	Asp	Met	Lys	Gly	His	Phe	Ser	Ile	Ser	Ile	515	520	525	
Pro	Val	Lys	Ser	Asp	Ile	Ala	Pro	Val	Ala	Arg	Leu	Leu	Ile	Tyr	Ala	530	535	540	
Val	Leu	Pro	Thr	Gly	Asp	Val	Ile	Gly	Asp	Ser	Ala	Lys	Tyr	Asp	Val	545	550	555	560
Glu	Asn	Cys	Leu	Ala	Asn	Lys	Val	Asp	Leu	Ser	Phe	Ser	Pro	Ser	Gln	565	570	575	
Ser	Leu	Pro	Ala	Ser	His	Ala	His	Leu	Arg	Val	Thr	Ala	Ala	Pro	Gln	580	585	590	
Ser	Val	Cys	Ala	Leu	Arg	Ala	Val	Asp	Gln	Ser	Val	Leu	Leu	Met	Lys	595	600	605	

Pro Asp Ala Glu Leu Ser Ala Ser Ser Val Tyr Asn Leu Leu Pro Glu
 610 615 620
 Lys Asp Leu Thr Gly Phe Pro Gly Pro Leu Asn Asp Gln Asp Asp Glu
 625 630 635 640
 Asp Cys Ile Asn Arg His Asn Val Tyr Ile Asn Gly Ile Thr Tyr Thr
 645 650 655
 Pro Val Ser Ser Thr Asn Glu Lys Asp Met Tyr Ser Phe Leu Glu Asp
 660 665 670
 Met Gly Leu Lys Ala Phe Thr Asn Ser Lys Ile Arg Lys Pro Lys Met
 675 680 685
 Cys Pro Gln Leu Gln Gln Tyr Glu Met His Gly Pro Glu Gly Leu Arg
 690 695 700
 Val Gly Phe Tyr Glu Ser Asp Val Met Gly Arg Gly His Ala Arg Leu
 705 710 715 720
 Val His Val Glu Glu Pro His Thr Glu Thr Val Arg Lys Tyr Phe Pro
 725 730 735
 Glu Thr Trp Ile Trp Asp Leu Val Val Val Asn Ser Ala Gly Val Ala
 740 745 750
 Glu Val Gly Val Thr Val Pro Asp Thr Ile Thr Glu Trp Lys Ala Gly
 755 760 765
 Ala Phe Cys Leu Ser Glu Asp Ala Gly Leu Gly Ile Ser Ser Thr Ala
 770 775 780
 Ser Leu Arg Ala Phe Gln Pro Phe Phe Val Glu Leu Thr Met Pro Tyr
 785 790 795 800
 Ser Val Ile Arg Gly Glu Ala Phe Thr Leu Lys Ala Thr Val Leu Asn
 805 810 815
 Tyr Leu Pro Lys Cys Ile Arg Val Ser Val Gln Leu Glu Ala Ser Pro
 820 825 830
 Ala Phe Leu Ala Val Pro Val Glu Lys Glu Gln Ala Pro His Cys Ile
 835 840 845
 Cys Ala Asn Gly Arg Gln Thr Val Ser Trp Ala Val Thr Pro Lys Ser
 850 855 860
 Leu Gly Asn Val Asn Phe Thr Val Ser Ala Glu Ala Leu Glu Ser Gln
 865 870 875 880
 Glu Leu Cys Gly Thr Glu Val Pro Ser Val Pro Glu His Gly Arg Lys
 885 890 895
 Asp Thr Val Ile Lys Pro Leu Leu Val Glu Pro Glu Gly Leu Glu Lys
 900 905 910
 Glu Thr Thr Phe Asn Ser Leu Leu Cys Pro Ser Gly Gly Glu Val Ser
 915 920 925
 Glu Glu Leu Ser Leu Lys Leu Pro Pro Asn Val Val Glu Glu Ser Ala
 930 935 940
 Arg Ala Ser Val Ser Val Leu Gly Asp Ile Leu Gly Ser Ala Met Gln
 945 950 955 960
 Asn Thr Gln Asn Leu Leu Gln Met Pro Tyr Gly Cys Gly Glu Gln Asn
 965 970 975
 Met Val Leu Phe Ala Pro Asn Ile Tyr Val Leu Asp Tyr Leu Asn Glu
 980 985 990
 Thr Gln Gln Leu Thr Pro Glu Val Lys Ser Lys Ala Ile Gly Tyr Leu
 995 1000 1005
 Asn Thr Gly Tyr Gln Arg Gln Leu Asn Tyr Lys His Tyr Asp Gly Ser
 1010 1015 1020
 Tyr Ser Thr Phe Gly Glu Arg Tyr Gly Arg Asn Gln Gly Asn Thr Trp
 1025 1030 1035 1040
 Leu Thr Ala Phe Val Leu Lys Thr Phe Ala Gln Ala Arg Ala Tyr Ile
 1045 1050 1055
 Phe Ile Asp Glu Ala His Ile Thr Gln Ala Leu Ile Trp Leu Ser Gln
 1060 1065 1070

Arg Gln Lys Asp Asn Gly Cys Phe Arg Ser Ser Gly Ser Leu Leu Asn
 1075 1080 1085
 Asn Ala Ile Lys Gly Gly Val Glu Asp Glu Val Thr Leu Ser Ala Tyr
 1090 1095 1100
 Ile Thr Ile Ala Leu Leu Glu Ile Pro Leu Thr Val Thr His Pro Val
 1105 1110 1115 1120
 Val Arg Asn Ala Leu Phe Cys Leu Glu Ser Ala Trp Lys Thr Ala Gln
 1125 1130 1135
 Glu Gly Asp His Gly Ser His Val Tyr Thr Lys Ala Leu Leu Ala Tyr
 1140 1145 1150
 Ala Phe Ala Leu Ala Gly Asn Gln Asp Lys Arg Lys Glu Val Leu Lys
 1155 1160 1165
 Ser Leu Asn Glu Glu Ala Val Lys Lys Asp Asn Ser Val His Trp Glu
 1170 1175 1180
 Arg Pro Gln Lys Pro Lys Ala Pro Val Gly His Phe Tyr Glu Pro Gln
 1185 1190 1195 1200
 Ala Pro Ser Ala Glu Val Glu Met Thr Ser Tyr Val Leu Leu Ala Tyr
 1205 1210 1215
 Leu Thr Ala Gln Pro Ala Pro Thr Ser Glu Asp Leu Thr Ser Ala Thr
 1220 1225 1230
 Asn Ile Val Lys Trp Ile Thr Lys Gln Gln Asn Ala Gln Gly Gly Phe
 1235 1240 1245
 Ser Ser Thr Gln Asp Thr Val Val Ala Leu His Ala Leu Ser Lys Tyr
 1250 1255 1260
 Gly Ala Ala Thr Phe Thr Arg Thr Gly Lys Ala Ala Gln Val Thr Ile
 1265 1270 1275 1280
 Gln Ser Ser Gly Thr Phe Ser Ser Lys Phe Gln Val Asp Asn Asn Asn
 1285 1290 1295
 Arg Leu Leu Leu Gln Gln Val Ser Leu Pro Glu Leu Pro Gly Glu Tyr
 1300 1305 1310
 Ser Met Lys Val Thr Gly Glu Gly Cys Val Tyr Leu Gln Thr Ser Leu
 1315 1320 1325
 Lys Tyr Asn Ile Leu Pro Glu Lys Glu Glu Phe Pro Phe Ala Leu Gly
 1330 1335 1340
 Val Gln Thr Leu Pro Gln Thr Cys Asp Glu Pro Lys Ala His Thr Ser
 1345 1350 1355 1360
 Phe Gln Ile Ser Leu Ser Val Ser Tyr Thr Gly Ser Arg Ser Ala Ser
 1365 1370 1375
 Asn Met Ala Ile Val Asp Val Lys Met Val Ser Gly Phe Ile Pro Leu
 1380 1385 1390
 Lys Pro Thr Val Lys Met Leu Glu Arg Ser Asn His Val Ser Arg Thr
 1395 1400 1405
 Glu Val Ser Ser Asn His Val Leu Ile Tyr Leu Asp Lys Val Ser Asn
 1410 1415 1420
 Gln Thr Leu Ser Leu Phe Phe Thr Val Leu Gln Asp Val Pro Val Arg
 1425 1430 1435 1440
 Asp Leu Lys Pro Ala Ile Val Lys Val Tyr Asp Tyr Tyr Glu Thr Asp
 1445 1450 1455
 Glu Phe Ala Ile Ala Glu Tyr Asn Ala Pro Cys Ser Lys Asp Leu Gly
 1460 1465 1470
 Asn Ala